

05-90
1009

#2

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/940,063

DATE: 10/01/2001

TIME: 13:58:13

Input Set : N:\Crf3\RULE60\09940063.txt

Output Set: N:\CRF3\10012001\I940063.raw

#1 2

4 <110> APPLICANT: Briskin, Michael J.
5 Murphy, Kristine E.
6 Wilbanks, Alyson M.
7 Wu, Lijun
9 <120> TITLE OF INVENTION: Novel Antibodies and Ligands for "Bonzo"
10 Chemokine Receptor
12 <130> FILE REFERENCE: 1855.1070-000
14 <140> CURRENT APPLICATION NUMBER: 09/940,063
15 <141> CURRENT FILING DATE: 2001-08-27
17 <150> PRIOR APPLICATION NUMBER: 09/449,437
18 <151> PRIOR FILING DATE: 1999-11-24
20 <160> NUMBER OF SEQ ID NOS: 18
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 1029
26 <212> TYPE: DNA
27 <213> ORGANISM: Homo sapiens
29 <400> SEQUENCE: 1

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31 gaggagcattc aagacttcct gcagttcagc aagggtcttc tgccctgcat gtacctggtg 120
32 gtgtttgtct gtggtctggt ggggaactct ctggtgctgg tcatatccat cttctaccat 180
33 aagttgcaga gcctgacgga tgtgttcctg gtgaacctac ccctggctga cctggtgttt 240
34 gtctgcactc tgcccttctg ggcctatgca ggcattccat aatgggtgtt tggccaggtc 300
35 atgtgcaaga gcctactggg catctacact attaatctct acacgtccat gctcatcctc 360
36 acctgcatca ctgtggatcg ttctattgta gtggttaagg ccaccaaggc ctacaaccag 420
37 caagccaaga ggatgacctg gggcaaggtc accagcttgc tcatctgggt gatatccctg 480
38 ctgggttctc tgccccaat tatctatggc aatgtcttta atctcgacaa gctcatatgt 540
39 gggttaccatg acgaggcaat ttccactgtg gttcttgcca cccagatgac actgggggttc 600
40 ttcttgccac tgetcaccat gattgtctgc tattcagtca taatcaaaac actgcttcat 660
41 gctggaggct tccagaagca cagatctcta aagatcatct tcttggtgat ggctgtgttc 720
42 ctgctgaccc agatgccctt caacctcatg aagttcatcc gcagcacaca ctgggaatac 780
43 tatgccatga ccagctttca ctacaccatc atggtgacag aggccatcgc atacctgagg 840
44 gcctgcctta accctgtgct ctatgccttt gtcagcctga agtttcgaaa gaacttctgg 900
45 aaacttgtga aggacattgg ttgcctccct taccttgagg tctcacatca atggaaatct 960
46 tctgaggaca attccaagac tttttctgcc tcccacaatg tggaggccac cagcatgttc 1020
47 cagttatag 1029

49 <210> SEQ ID NO: 2
50 <211> LENGTH: 342
51 <212> TYPE: PRT
52 <213> ORGANISM: Homo sapiens
54 <400> SEQUENCE: 2

55 Met Ala Glu His Asp Tyr His Glu Asp Tyr Gly Phe Ser Ser Phe Asn
56 1 5 10 15
57 Asp Ser Ser Gln Glu Glu His Gln Asp Phe Leu Gln Phe Ser Lys Val
58 20 25 30
59 Phe Leu Pro Cys Met Tyr Leu Val Val Phe Val Cys Gly Leu Val Gly
60 35 40 45

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61 Asn Ser Leu Val Leu Val Ile Ser Ile Phe Tyr His Lys Leu Gln Ser
62      50                      55                      60
63 Leu Thr Asp Val Phe Leu Val Asn Leu Pro Leu Ala Asp Leu Val Phe
64 65      70                      75                      80
65 Val Cys Thr Leu Pro Phe Trp Ala Tyr Ala Gly Ile His Glu Trp Val
66      85                      90                      95
67 Phe Gly Gln Val Met Cys Lys Ser Leu Leu Gly Ile Tyr Thr Ile Asn
68      100                     105                     110
69 Phe Tyr Thr Ser Met Leu Ile Leu Thr Cys Ile Thr Val Asp Arg Phe
70      115                     120                     125
71 Ile Val Val Val Lys Ala Thr Lys Ala Tyr Asn Gln Gln Ala Lys Arg
72      130                     135                     140
73 Met Thr Trp Gly Lys Val Thr Ser Leu Leu Ile Trp Val Ile Ser Leu
74 145      150                     155                     160
75 Leu Val Ser Leu Pro Gln Ile Ile Tyr Gly Asn Val Phe Asn Leu Asp
76      165                     170                     175
77 Lys Leu Ile Cys Gly Tyr His Asp Glu Ala Ile Ser Thr Val Val Leu
78      180                     185                     190
79 Ala Thr Gln Met Thr Leu Gly Phe Phe Leu Pro Leu Leu Thr Met Ile
80      195                     200                     205
81 Val Cys Tyr Ser Val Ile Ile Lys Thr Leu Leu His Ala Gly Gly Phe
82      210                     215                     220
83 Gln Lys His Arg Ser Leu Lys Ile Ile Phe Leu Val Met Ala Val Phe
84 225      230                     235                     240
85 Leu Leu Thr Gln Met Pro Phe Asn Leu Met Lys Phe Ile Arg Ser Thr
86      245                     250                     255
87 His Trp Glu Tyr Tyr Ala Met Thr Ser Phe His Tyr Thr Ile Met Val
88      260                     265                     270
89 Thr Glu Ala Ile Ala Tyr Leu Arg Ala Cys Leu Asn Pro Val Leu Tyr
90      275                     280                     285
91 Ala Phe Val Ser Leu Lys Phe Arg Lys Asn Phe Trp Lys Leu Val Lys
92      290                     295                     300
93 Asp Ile Gly Cys Leu Pro Tyr Leu Gly Val Ser His Gln Trp Lys Ser
94 305      310                     315                     320
95 Ser Glu Asp Asn Ser Lys Thr Phe Ser Ala Ser His Asn Val Glu Ala
96      325                     330                     335
97 Thr Ser Met Phe Gln Leu
98      340
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102 <211> LENGTH: 1763
103 <212> TYPE: DNA
104 <213> ORGANISM: Homo sapiens
106 <400> SEQUENCE: 3
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108 ttctgctcct gctggtgtac ctgactcagc caggcaatgg caacgagggc agcgtcactg 120
109 gaagttgtta ttgtggtaaa agaatttctt ccgactcccc gccatcgggt cagttcatga 180
110 atcgtctccg gaaacacctg agagcttacc atcgggtgtct atactacacg aggttccagc 240
111 tccttttctg gagcgtgtgt ggaggcaaca aggacccatg ggttcaggaa ttgatgagct 300
112 gtcttgatct caaagaatgt ggacatgctt actcggggat tgtggcccac cagaagcatt 360

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113 tacttcctac cagcccccca atttctcagg cctcagaggg ggcattctca gatataccaca 420
114 cccctgccca gatgctcctg tccaccttgc agtccactca gcgccccacc ctcccagtag 480
115 gatcactgtc ctoggacaaa gagctcactc gtcccaatga aaccaccatt cacactgcgg 540
116 gccacagtct ggcagttggg cctgaggctg gggagaacca gaagcagccg gaaaaaaatg 600
117 ctggtcccac agccaggaca tcagccacag tgccggctct gtgcctcctg gccatcatct 660
118 tcactcctac cgcagccctt tcctatgtgc tgtgcaagag gaggaggggg cagtcaccgc 720
119 agtcctctcc agatctgccg gttcattata tacctgtggc acctgactct aatacctgag 780
120 ccaagaatgg aagcttgtga ggagacggac tctatgttgc ccaggctgtt atggaactcc 840
121 tgagtcaagt gatcctccca ccttggcctc tgaagggtgc aggattatag gcgtcaccta 900
122 ccacatccag cctacacgta tttgttaata tctaacatag gactaaccag ccactgccct 960
123 ctcttagggc cctcatttaa aaacggttat actataaaat ctgcttttca cactgggtga 1020
124 taataacttg gacaaattct atgtgtatgt tgttttgttt tgctttgctt tgttttgaga 1080
125 cggagtctcg ctctgtcctc caggctggag tgcagtggca tgatctcggc tcaactgcaac 1140
126 ccccatctcc caggttcaag cgattctcct gcctcctcct gagtagctgg gactacaggt 1200
127 gctcaccacc acaccggct aattttttgt attttttagta gagaccgggg ttccaccatg 1260
128 ttgaccaggc tggctctgaa ctctgacct ggtgatctgc ccaccaggc ctcccaaagt 1320
129 gctgggatta aaggtgtgag ccaccatgcc tggccctatg tgtgtttttt aactactaaa 1380
130 aattattttt gtaatgattg agtcttcttt atggaacaaa ctggcctcag cccttgccgc 1440
131 ctactgtgta ttctggctt cattttttgc tgatggttcc cctcgtccc aaatctctct 1500
132 cccagtacac cagttgttcc tccccacct cagccctctc ctgcatctc ctgtaccgc 1560
133 aacgaaggcc tgggttttcc caccctcct ccttagcagg tgccgtgctg ggacaccata 1620
134 cgggttggtt tcacctctc agtcccttgc ctaccocagt gagagtctga tcttgttttt 1680
135 attgttattg cttttattat tattgctttt attatcatta aaactctagt tcttgttttg 1740
136 tctctccgaa aaaaaaaaaa aaa 1763

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138 <210> SEQ ID NO: 4

139 <211> LENGTH: 254

140 <212> TYPE: PRT

141 <213> ORGANISM: Homo sapiens

143 <400> SEQUENCE: 4

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146 Leu Leu Leu Leu Val Tyr Leu Thr Gln Pro Gly Asn Gly Asn Glu Gly
147 20 25 30
148 Ser Val Thr Gly Ser Cys Tyr Cys Gly Lys Arg Ile Ser Ser Asp Ser
149 35 40 45
150 Pro Pro Ser Val Gln Phe Met Asn Arg Leu Arg Lys His Leu Arg Ala
151 50 55 60
152 Tyr His Arg Cys Leu Tyr Tyr Thr Arg Phe Gln Leu Leu Ser Trp Ser
153 65 70 75 80
154 Val Cys Gly Gly Asn Lys Asp Pro Trp Val Gln Glu Leu Met Ser Cys
155 85 90 95
156 Leu Asp Leu Lys Glu Cys Gly His Ala Tyr Ser Gly Ile Val Ala His
157 100 105 110
158 Gln Lys His Leu Leu Pro Thr Ser Pro Pro Ile Ser Gln Ala Ser Glu
159 115 120 125
160 Gly Ala Ser Ser Asp Ile His Thr Pro Ala Gln Met Leu Leu Ser Thr
161 130 135 140
162 Leu Gln Ser Thr Gln Arg Pro Thr Leu Pro Val Gly Ser Leu Ser Ser
163 145 150 155 160

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164 Asp Lys Glu Leu Thr Arg Pro Asn Glu Thr Thr Ile His Thr Ala Gly
165                               165                               170                               175
166 His Ser Leu Ala Val Gly Pro Glu Ala Gly Glu Asn Gln Lys Gln Pro
167                               180                               185                               190
168 Glu Lys Asn Ala Gly Pro Thr Ala Arg Thr Ser Ala Thr Val Pro Val
169                               195                               200                               205
170 Leu Cys Leu Leu Ala Ile Ile Phe Ile Leu Thr Ala Ala Pro Ser Tyr
171                               210                               215                               220
172 Val Leu Cys Lys Arg Arg Arg Gly Gln Ser Pro Gln Ser Ser Pro Asp
173 225                               230                               235                               240
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175                               245                               250
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179 <211> LENGTH: 2309
180 <212> TYPE: DNA
181 <213> ORGANISM: Homo sapiens
183 <400> SEQUENCE: 5
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185 gcccggatcg gggaagtgaa agtgccctcg aggaggagg ccggtccggc agtgacgccg 120
186 cctcacaggt cggcggacgg gccaggcggg cggcctcctg aaccgaaccg aatcggtcc 180
187 tcgggcccgc gtccctccgc cctcctcgc ccgcgcgcgg agttttcttt cggtttcttc 240
188 caagattcct ggccttccct cgaaggagcc gggcccagtg cgggggcgca gggcgcgga 300
189 gctccacctc ctgggttttc cctgcgtcca gaggtggca tggcgcgggc cgagtactga 360
190 ggcacacggtc ggggcacagc agggccgggt ggtgcagctg gctcgcgcct cctctccggc 420
191 cgcgctctcc tcgggtcccc ggcgaaagcc attgagacac cagctggacg tcacgcgccg 480
192 gagcatgtct gggagtcaga gcgaggtggc tccatccccg cagagtccgc ggagccccga 540
193 gatgggacgg gacttgccgc cgggtcccg cgtgctcctg ctctgtcttc tgctcctgct 600
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195 tggtaaaaga atttcttccg actccccgcc atcggttcag ttcataaata gtctccggaa 720
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197 cgtgtgtgga ggcaacaagg acctatgggt tcaggaattg atgagctgtc ttgatctcaa 840
198 agaattgtgga catgcttact cggggattgt ggcccaccag aagcatttac ttctaccag 900
199 ccccccactc tctcaggcct cagagggggc atcttcagat atccacaccc ctgccagat 960
200 gctcctgtcc accttgacgt ccactcagcg cccaccctc ccagtaggat cactgtcctc 1020
201 ggacaaagag ctactcgtc ccaatgaaac caccattcac actgcggggc acagtctggc 1080
202 agttgggcct gaggctggg agaaccagaa gcagccggaa aaaaatgctg gtcccacagc 1140
203 caggacatca gccacagtgc cggctcctgt cctcctggcc atcatcttca tctcaccgc 1200
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207 cctcccacct tggcctctga aggtgcgagg attataggcg tcacctacca catccagcct 1440
208 acacgtatct gttaatatct aacataggac taaccagcca ctgccctctc ttaggccctc 1500
209 catttaaaaa cggttatact ataaaatctg cttttcacac tgggtgataa taacttgga 1560
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211 tgtcatccag gctggagtgc agtggcatga tctcggtcca ctgcaacccc catctcccag 1680
212 gttcaagcga ttctcctgcc tcctcctaag tagctgggac tacaggtgct caccaccaca 1740
213 cccggctaata tttttgtatt tttagtagag acggggtttc accatgttga ccaggctggt 1800
214 ctcgaaactcc tgacctggtg atctgccac ccaggcctcc caaagtgctg ggattaaagg 1860
215 tgtgagccac catgcctggc cctatgtgtg ttttttaact actaaaaatt atttttgtaa 1920

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216 tgattgagtc ttctttatgg aaacaactgg cctcagccct tgcgccctta ctgtgattcc 1980
217 tggcttcatt ttttgctgat gggtcccccct cgtcccaaat ctctctccca gtacaccagt 2040
218 tgttcctccc ccaacctcagc cctctcctgc atcctcctgt acccgcaacg aaggcctggg 2100
219 ctttcccacc ctccctcctt agcagggtgcc gtgctgggac accatacggg ttggtttcac 2160
220 ctccctcagtc ccttgcttac ccagtgaga gtctgatctt gtttttattg ttattgcttt 2220
221 tattattatt gcttttatta tcattaaaac tctagttctt gttttgtctc tcaaaaaaaaa 2280
222 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 2309

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224 <210> SEQ ID NO: 6

225 <211> LENGTH: 254

226 <212> TYPE: PRT

227 <213> ORGANISM: Homo sapiens

229 <400> SEQUENCE: 6

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233 20 25 30
234 Ser Val Thr Gly Ser Cys Tyr Cys Gly Lys Arg Ile Ser Ser Asp Ser
235 35 40 45
236 Pro Pro Ser Val Gln Phe Met Asn Arg Leu Arg Lys His Leu Arg Ala
237 50 55 60
238 Tyr His Arg Cys Leu Tyr Tyr Thr Arg Phe Gln Leu Leu Ser Trp Ser
239 65 70 75 80
240 Val Cys Gly Gly Asn Lys Asp Pro Trp Val Gln Glu Leu Met Ser Cys
241 85 90 95
242 Leu Asp Leu Lys Glu Cys Gly His Ala Tyr Ser Gly Ile Val Ala His
243 100 105 110
244 Gln Lys His Leu Leu Pro Thr Ser Pro Pro Thr Ser Gln Ala Ser Glu
245 115 120 125
246 Gly Ala Ser Ser Asp Ile His Thr Pro Ala Gln Met Leu Leu Ser Thr
247 130 135 140
248 Leu Gln Ser Thr Gln Arg Pro Thr Leu Pro Val Gly Ser Leu Ser Ser
249 145 150 155 160
250 Asp Lys Glu Leu Thr Arg Pro Asn Glu Thr Thr Ile His Thr Ala Gly
251 165 170 175
252 His Ser Leu Ala Val Gly Pro Glu Ala Gly Glu Asn Gln Lys Gln Pro
253 180 185 190
254 Glu Lys Asn Ala Gly Pro Thr Ala Arg Thr Ser Ala Thr Val Pro Val
255 195 200 205
256 Leu Cys Leu Leu Ala Ile Ile Phe Ile Leu Thr Ala Ala Leu Ser Tyr
257 210 215 220
258 Val Leu Cys Lys Arg Arg Arg Gly Gln Ser Pro Gln Ser Ser Pro Asp
259 225 230 235 240
260 Leu Pro Val His Tyr Ile Pro Val Ala Pro Asp Ser Asn Thr
261 245 250

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264 <210> SEQ ID NO: 7

265 <211> LENGTH: 439

266 <212> TYPE: DNA

267 <213> ORGANISM: Homo sapiens

269 <400> SEQUENCE: 7

VERIFICATION SUMMARY

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